

PCT/US03/28654 .11032005

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SEQUENCE LISTING

<110> Behrens, Sven-Erik
Isken, Olaf
Grassmann, Claus W.
Sarisky, Robert T.

<120> A Set Of Ubiquitous Cellular Proteins
Involved in Viral Life Cycle

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 115 120 125
 Ala Asp Asn Leu Ala Ile Gln Leu Ala Ala Val Thr Glu Asp Lys Tyr
 130 135 140
 Glu Ile Leu Gln Ser Val Asp Asp Ala Ala Ile Val Ile Lys Asn Thr
 145 150 155 160
 Lys Glu Pro Pro Leu Ser Leu Thr Ile His Leu Thr Ser Pro Val Val
 165 170 175
 Arg Glu Glu Met Glu Lys Val Leu Ala Gly Glu Thr Leu Ser Val Asn
 180 185 190
 Asp Pro Pro Asp Val Leu Asp Arg Gln Lys Cys Leu Ala Ala Leu Ala
 195 200 205
 Ser Leu Arg His Ala Lys Trp Phe Gln Ala Arg Ala Asn Gly Leu Lys
 210 215 220
 Ser Cys Val Ile Val Ile Arg Val Leu Arg Asp Leu Cys Thr Arg Val
 225 230 235 240
 Pro Thr Trp Gly Pro Leu Arg Gly Trp Pro Leu Glu Leu Leu Cys Glu
 245 250 255

SEQLIST.TXT

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Lys Ser Ile Gly Thr Ala Asn Arg Pro Met Gly Ala Gly Glu Ala Leu
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Arg Arg Val Leu Glu Cys Leu Ala Ser Gly Ile Val Met Pro Asp Gly
      275      280      285
Ser Gly Ile Tyr Asp Pro Cys Glu Lys Glu Ala Thr Asp Ala Ile Gly
      290      295      300
His Leu Asp Arg Gln Gln Arg Glu Asp Ile Thr Gln Ser Ala Gln His
      305      310      315      320
Ala Leu Arg Leu Ala Ala Phe Gly Gln Leu His Lys Val Leu Gly Met
      325      330      335
Asp Pro Leu Pro Ser Lys Met Pro Lys Lys Pro Lys Asn Glu Asn Pro
      340      345      350
Val Asp Tyr Thr Val Gln Ile Pro Pro Ser Thr Thr Tyr Ala Ile Thr
      355      360      365
Pro Met Lys Arg Pro Met Glu Glu Asp Gly Glu Glu Lys Ser Pro Ser
      370      375      380
Lys Lys Lys Lys Lys Ile Gln Lys Lys Glu Glu Lys Ala Glu Pro Pro
      385      390      395      400
Gln Ala Met Asn Ala Leu Met Arg Leu Asn Gln Leu Lys Pro Gly Leu
      405      410      415
Gln Tyr Lys Leu Val Ser Gln Thr Gly Pro Val His Ala Pro Ile Phe
      420      425      430
Thr Met Ser Val Glu Val Asp Gly Asn Ser Phe Glu Ala Ser Gly Pro
      435      440      445
Ser Lys Lys Thr Ala Lys Leu His Val Ala Val Lys Val Leu Gln Asp
      450      455      460
Met Gly Leu Pro Thr Gly Ala Glu Gly Arg Asp Ser Ser Lys Gly Glu
      465      470      475      480
Asp Ser Ala Glu Glu Thr Glu Ala Lys Pro Ala Val Val Ala Pro Ala
      485      490      495
Pro Val Val Glu Ala Val Ser Thr Pro Ser Ala Ala Phe Pro Ser Asp
      500      505      510
Ala Thr Ala Glu Gln Gly Pro Ile Leu Thr Lys His Gly Lys Asn Pro
      515      520      525
Val Met Glu Leu Asn Glu Lys Arg Arg Gly Leu Lys Tyr Glu Leu Ile
      530      535      540
Ser Glu Thr Gly Gly Ser His Asp Lys Arg Phe Val Met Glu Val Glu
      545      550      555      560
Val Asp Gly Gln Lys Phe Gln Gly Ala Gly Ser Asn Lys Lys Val Ala
      565      570      575
Lys Ala Tyr Ala Ala Leu Ala Ala Leu Glu Lys Leu Phe Pro Asp Thr
      580      585      590
Pro Leu Ala Leu Asp Ala Asn Lys Lys Lys Arg Ala Pro Val Pro Val
      595      600      605
Arg Gly Gly Pro Lys Phe Ala Ala Lys Pro His Asn Pro Gly Phe Gly
      610      615      620
Met Gly Gly Pro Met His Asn Glu Val Pro Pro Pro Pro Asn Leu Arg
      625      630      635      640
Gly Arg Gly Arg Gly Ser Ile Arg Gly Arg Gly Arg Gly Arg Gly
      645      650      655
Phe Gly Gly Ala Asn His Gly Gly Tyr Met Asn Ala Gly Ala Gly Tyr
      660      665      670
Gly Ser Tyr Gly Tyr Gly Gly Asn Ser Ala Thr Ala Gly Tyr Ser Asp
      675      680      685
Phe Phe Thr Asp Cys Tyr Gly Tyr His Asp Phe Gly Ser Ser
      690      695      700

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<210> 6

<211> 2107

<212> DNA

<213> Homo sapien

<400> 6

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gcgctcaaa ctgtgtccga ctggatagac gacgaggaaa agggtagcag cgagcaggca 180
gagtccgata acatggatgt gccccagag gacgacagta aagaaggggc tggggaacag 240

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SEQLIST.TXT

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acaaccgccc tcctggacaa ggtggccgac aacctggcca tccagcttgc tgctgtaaca 420
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cccacctggg gtccccctcg aggctggcct ctcgagctcc tgtgtgagaa atccattggc 780
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<210> 7
 <211> 406
 <212> PRT
 <213> Homo sapien

<400> 7
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 35 40 45
 Asp Glu Thr Ser Phe Ser Glu Ala Leu Leu Lys Arg Asn Gln Asp Leu
 50 55 60
 Ala Pro Asn Ser Ala Glu Gln Ala Ser Ile Leu Ser Leu Val Thr Lys
 65 70 75 80
 Ile Asn Asn Val Ile Asp Asn Leu Ile Val Ala Pro Gly Thr Phe Glu
 85 90 95
 Val Gln Ile Glu Glu Val Arg Gln Val Gly Ser Tyr Lys Lys Gly Thr
 100 105 110
 Met Thr Thr Gly His Asn Val Ala Asp Leu Val Val Ile Leu Lys Ile
 115 120 125
 Leu Pro Thr Leu Glu Ala Val Ala Ala Leu Gly Asn Lys Val Val Glu
 130 135 140
 Ser Leu Arg Ala Gln Asp Pro Ser Glu Val Leu Thr Met Leu Thr Asn
 145 150 155 160
 Glu Thr Gly Phe Glu Ile Ser Ser Ser Asp Ala Thr Val Lys Ile Leu
 165 170 175
 Ile Thr Thr Val Pro Pro Asn Leu Arg Lys Leu Asp Pro Glu Leu His
 180 185 190
 Leu Asp Ile Lys Val Leu Gln Ser Ala Leu Ala Ala Ile Arg His Ala
 195 200 205
 Arg Trp Phe Glu Glu Asn Ala Ser Gln Ser Thr Val Lys Val Leu Ile
 210 215 220
 Arg Leu Leu Lys Asp Leu Arg Ile Arg Phe Pro Gly Phe Glu Pro Leu

SEQLIST.TXT

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225          230          235          240
Thr Pro Trp Ile Leu Asp Leu Leu Gly His Tyr Ala Val Met Asn Asn
          245          250          255
Pro Thr Arg Gln Pro Leu Ala Leu Asn Val Ala Tyr Arg Arg Cys Leu
          260          265          270
Gln Ile Leu Ala Ala Gly Leu Phe Leu Pro Gly Ser Val Gly Ile Thr
          275          280          285
Asp Pro Cys Glu Ser Gly Asn Phe Arg Val His Thr Val Met Thr Leu
          290          295          300
Glu Gln Gln Asp Met Val Cys Tyr Thr Ala Gln Thr Leu Val Arg Ile
305          310          315          320
Leu Ser His Gly Gly Phe Arg Lys Ile Leu Gly Gln Glu Gly Asp Ala
          325          330          335
Ser Tyr Leu Ala Ser Glu Ile Ser Thr Trp Asp Gly Val Ile Val Thr
          340          345          350
Pro Ser Glu Lys Ala Tyr Glu Lys Pro Pro Glu Lys Lys Glu Gly Glu
          355          360          365
Glu Glu Glu Glu Asn Thr Glu Arg Thr Thr Ser Arg Arg Gly Arg Arg
          370          375          380
Lys His Gly Asn Ser Gly Val Thr Phe Pro Ser Leu Leu Phe Leu Pro
385          390          395          400
Lys Gly Lys Thr Gly Ala
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<210> 8
<211> 1221
<212> DNA
<213> Homo sapien

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aagggaaaga ctggagccta a
          1221

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